

## SEQUENCE LISTING

&lt;110&gt; ANDREEV, et al.

&lt;120&gt; METHODS FOR IDENTIFYING MODULATORS OF ACTIVE KIT TYROSINE KINASE RECEPTOR

&lt;130&gt; 30694/39618A

&lt;150&gt; US 60/526,930

&lt;151&gt; 2003-12-04

&lt;160&gt; 6

&lt;170&gt; PatentIn version 3.2

&lt;210&gt; 1

&lt;211&gt; 5084

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (22)..(2952)

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Gly Gly Thr Tyr Thr Phe Leu Val Ser Asn Ser Asp Val Ser Ala Ser	
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Val Thr Phe Asp Val Tyr Val Asn Thr Lys Pro Glu Ile Leu Thr Tyr	
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Asp Arg Leu Met Asn Gly Arg Leu Gln Cys Val Ala Ala Gly Phe Pro	
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Glu Pro Thr Ile Asp Trp Tyr Phe Cys Thr Gly Ala Glu Gln Arg Cys	
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Thr Val Pro Val Pro Pro Val Asp Val Gln Ile Gln Asn Ala Ser Val	
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Ser Pro Phe Gly Lys Leu Val Val Gln Ser Ser Ile Asp Ser Ser Val	
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Phe Arg His Asn Gly Thr Val Glu Cys Lys Ala Ser Asn Ala Val Gly	
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Lys Ser Ser Ala Phe Phe Asn Phe Ala Phe Lys Gly Asn Ser Lys Glu	
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Gln Ile Gln Pro His Thr Leu Phe Thr Pro Leu Leu Ile Gly Phe Val	
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Val Thr Ala Gly Leu Met Gly Ile Ile Val Met Val Leu Ala Tyr Lys	
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Leu	Ile	Lys	Ser	Asp	Ala	Ala	Met	Thr	Val	Ala	Val	Lys	Met	Leu	Lys	
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Pro	Ser	Ala	His	Leu	Thr	Glu	Arg	Glu	Ala	Leu	Met	Ser	Glu	Leu	Lys	
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Phe	Ser	Lys	Gln	Glu	Glu	Gln	Ala	Asp	Ala	Ala	Leu	Tyr	Lys	Asn	Leu	
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Leu	His	Ser	Lys	Glu	Ser	Ser	Cys	Asp	Ser	Ser	Asn	Glu	Tyr	Met	Asp	
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Arg	Ser	Ala	Arg	Ile	Asp	Ser	Tyr	Ile	Glu	Arg	Asp	Val	Thr	Pro	Ala	
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Ile	Met	Glu	Asp	Asp	Glu	Leu	Ala	Leu	Asp	Leu	Glu	Asp	Leu	Leu	Ser	
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Phe	Ser	Tyr	Gln	Val	Ala	Lys	Gly	Met	Ala	Phe	Leu	Ala	Ser	Lys	Asn	
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Thr Phe Lys Gln Val Val Gln Leu Ile Glu Lys Gln Ile Ser Asp Ser	
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Ser Lys His Ile Tyr Ser Asn Leu Ala Asn Cys Asn Pro Asn Pro Glu	
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Ser Thr Ser Ser Thr Gln Pro Leu Leu Val His Glu Asp Ala	
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Glu Ala Gly Asp Thr Ile Arg Leu Thr Cys Thr Asp Pro Ala Phe Val  
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Lys Trp Thr Phe Glu Ile Leu Asp Val Arg Ile Glu Asn Lys Gln Ser  
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Glu Trp Ile Arg Glu Lys Ala Glu Ala Thr His Thr Gly Lys Tyr Thr  
85 90 95

Cys Val Ser Gly Ser Gly Leu Arg Ser Ser Ile Tyr Val Phe Val Arg  
100 105 110

Asp Pro Ala Val Leu Phe Leu Val Gly Leu Pro Leu Phe Gly Lys Glu  
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Asp Asn Asp Ala Leu Val Arg Cys Pro Leu Thr Asp Pro Gln Val Ser  
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Asn Tyr Ser Leu Ile Glu Cys Asp Gly Lys Ser Leu Pro Thr Asp Leu  
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Lys Phe Val Pro Asn Pro Lys Ala Gly Ile Thr Ile Lys Asn Val Lys  
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Arg Ala Tyr His Arg Leu Cys Ile Arg Cys Ala Ala Gln Arg Glu Gly  
180 185 190



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 Glu Gly Asp Thr Phe Thr Val Ile Cys Thr Ile Lys Asp Val Ser Thr  
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 Ser Val Asp Ser Met Trp Ile Lys Leu Asn Pro Gln Pro Gln Ser Lys  
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Tyr Lys Asn Leu Leu His Ser Lys Glu Ser Ser Cys Asp Ser Ser Asn  
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Ser Val Gly Ser Ser Thr Ser Ser Thr Gln Pro Leu Leu Val His Glu  
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Asp Ala